

Working with Chromatogram

This chapter explains how to work efficiently with the *Sanger Reads Editor*.

- [Mapping Reads to Reference](#)
- [Alignment Statistics](#)
- [Alignment Appearance](#)
- [Overview and Show and Hide Chromatogram](#)
- [Sanger Reads Consensus](#)
 - [Export Chromatogram Consensus](#)
- [Navigation in Sanger Reads Alignment](#)
- [Editing Sanger Reads](#)
 - [Inserting Character](#)
 - [Replacing Character and Gap](#)
 - [Removing Character and Gap](#)
 - [Inserting Gap](#)
 - [Removing Gap at the Left](#)
 - [Removing All Columns of Gaps](#)
 - [Trimming Left End](#)
 - [Trimming Right End](#)
 - [Renaming Read](#)
 - [Removing Read](#)
 - [Undo and Redo](#)
- [Exporting Alignment without Chromatograms](#)
- [Sanger Reads Settings](#)